

In the Claims:

1-123. Canceled.

124. (Currently amended) An isolated nucleic acid comprising:

- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 387;
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 387, lacking its associated signal peptide;
- (c) ~~a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 387;~~
- (d) the nucleic acid sequence of SEQ ID NO: 386;
- (e)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 386;
- or
- (f)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203132.

125. (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 387.

126. (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 387, lacking its associated signal peptide.

127-128. Canceled.

129. (Previously presented) The isolated nucleic acid of Claim 124 comprising the nucleic acid sequence of SEQ ID NO: 386.

130. (Previously presented) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 386.

131. (Previously presented) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203132.
- 132-134. (Canceled)
135. (Previously presented) A vector comprising the nucleic acid of Claim 143 or 124.
136. (Previously presented) The vector of Claim 135, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
137. (Previously presented) An isolated host cell comprising the vector of Claim 135.
138. (Previously presented) The host cell of Claim 137, wherein said cell is a CHO cell, an *E. coli* or a yeast cell.
- 139-142. (Canceled)
143. (Currently amended) An isolated nucleic acid encoding a polypeptide having at least 99% sequence identity to:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 387;
 - (b) the amino acid sequence of the polypeptide of SEQ ID NO: 387, lacking its associated signal peptide;
 - (c) ~~a amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO: 387;~~
 - (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 386; or
 - (e)(d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203132; wherein said encoded polypeptide induces chondrocyte redifferentiation.